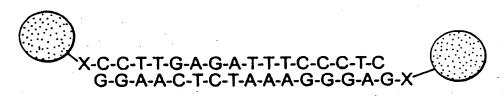
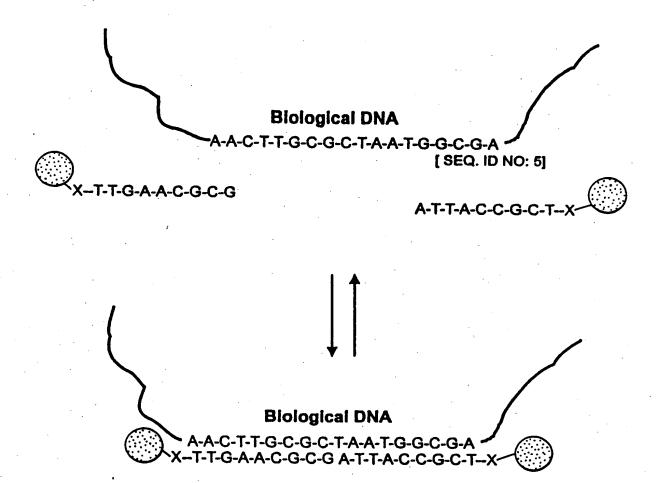
[SEQ. ID NO: 3]

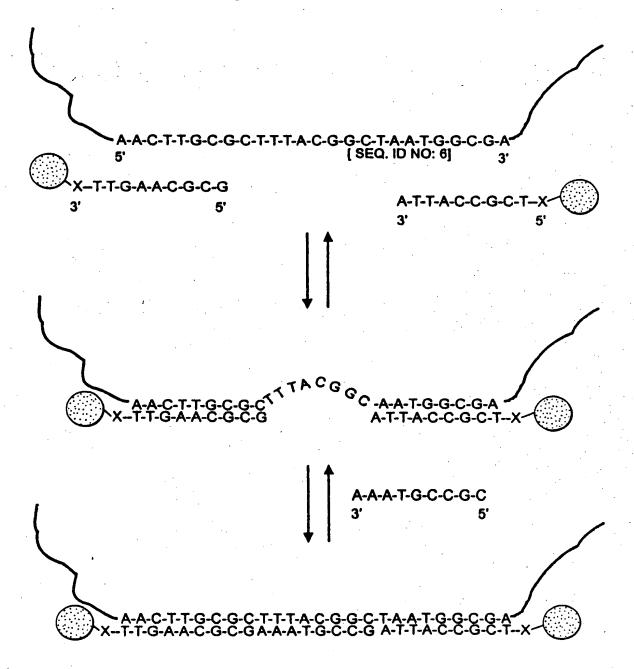
X-C-C-T-T-G-A-G-A-T-T-C-C-C-T-C
5'

G-G-A-A-C-T-C-T-A-A-A-G-G-G-A-G-X
[SEQ. ID NO: 4]









Precipitate (formed by further cross-linking)

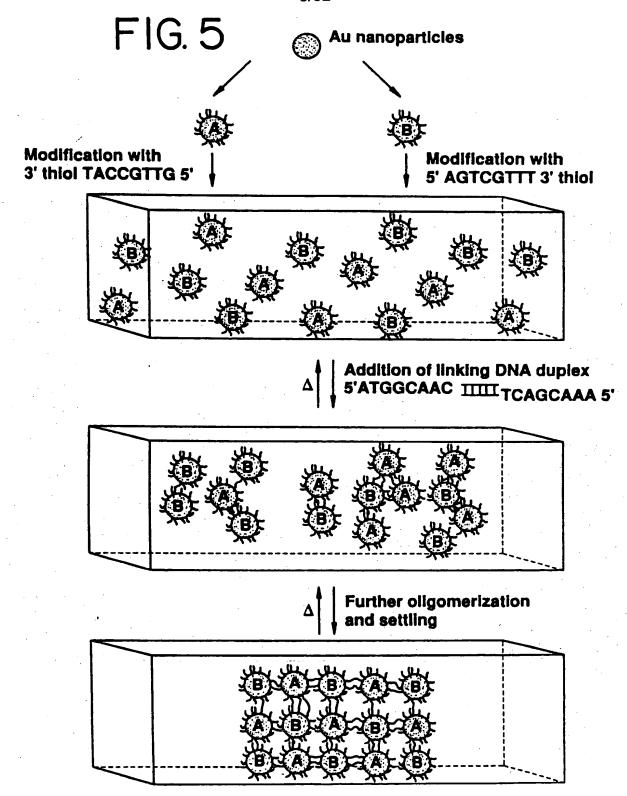
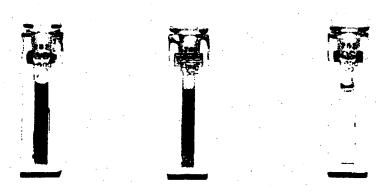
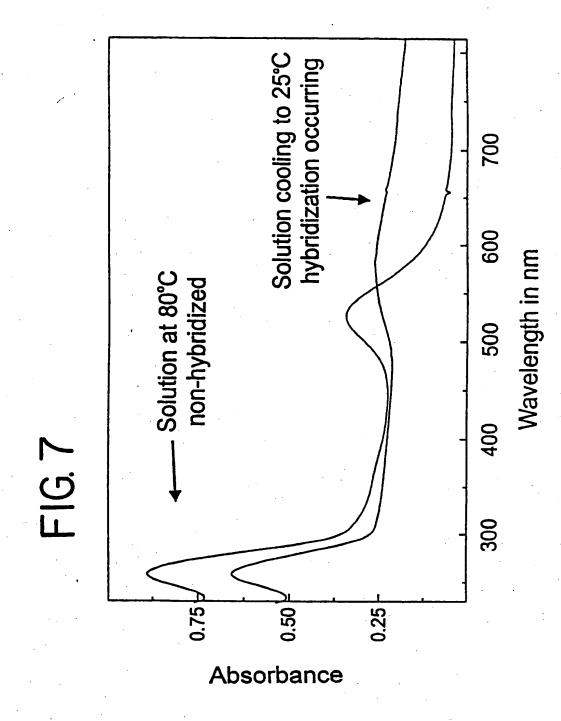


FIG. 6A FIG. 6B FIG. 6C





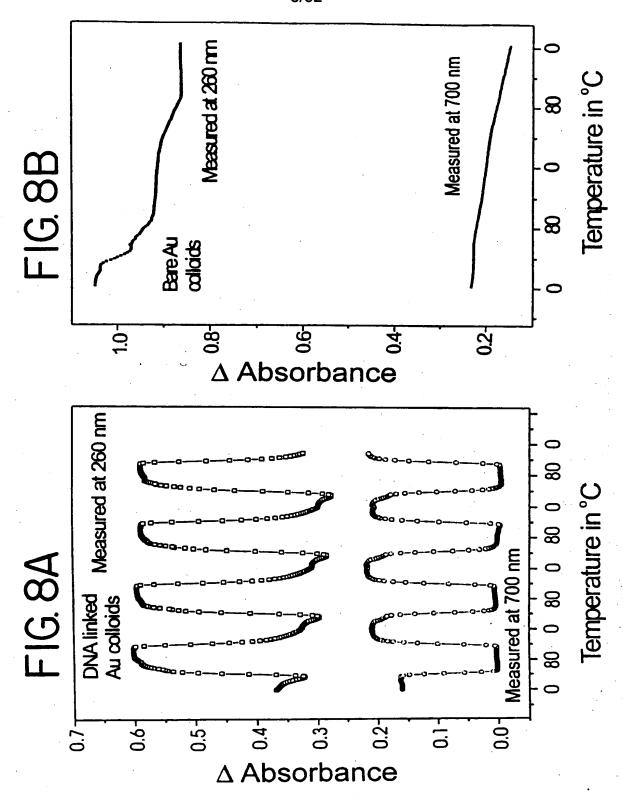


FIG. 9A



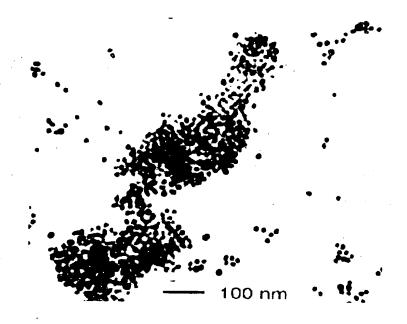
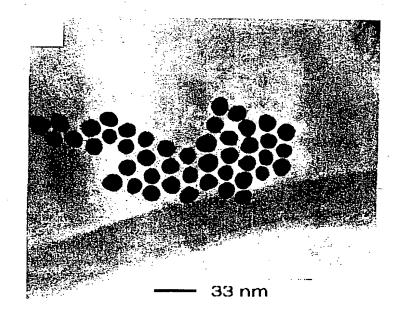


FIG. 9B

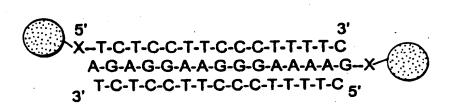


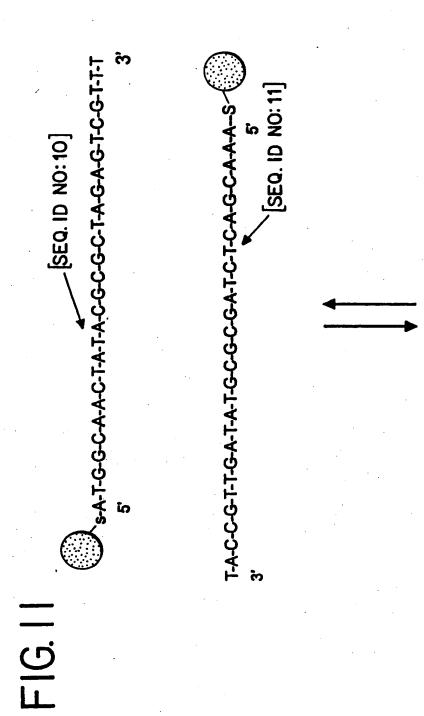
[SEQ ID NO: 7]

5'
3'
X--T-C-T-C-C-C-T-T-T-C
A-G-A-G-G-A-A-A-A-G-X
3'

[SEQ ID NO: 8]

3' T-C-T-C-C-T-T-C-C-C-T-T-T-T-C 5' [SEQ ID NO: 9]





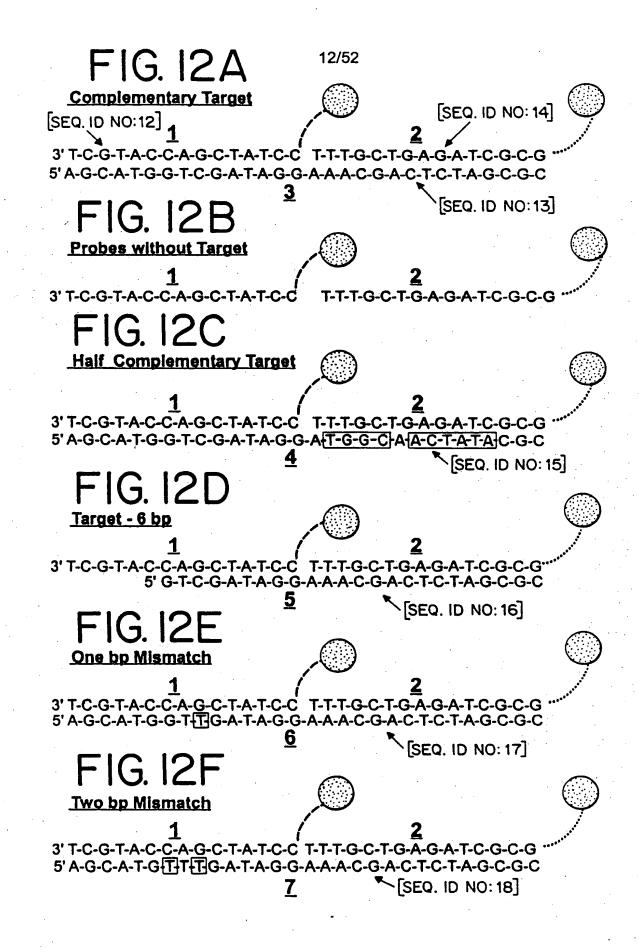


FIG. 13A

transparent substrate

Modified DNA chemisorbed onto solid substrate

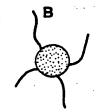
∫B'

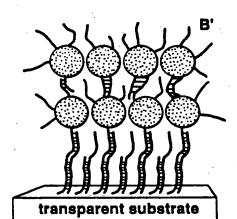
Analyte DNA

transparent substrate

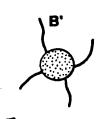
Analyte DNA hybridized onto substrate

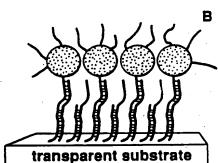
DNA modified colloids





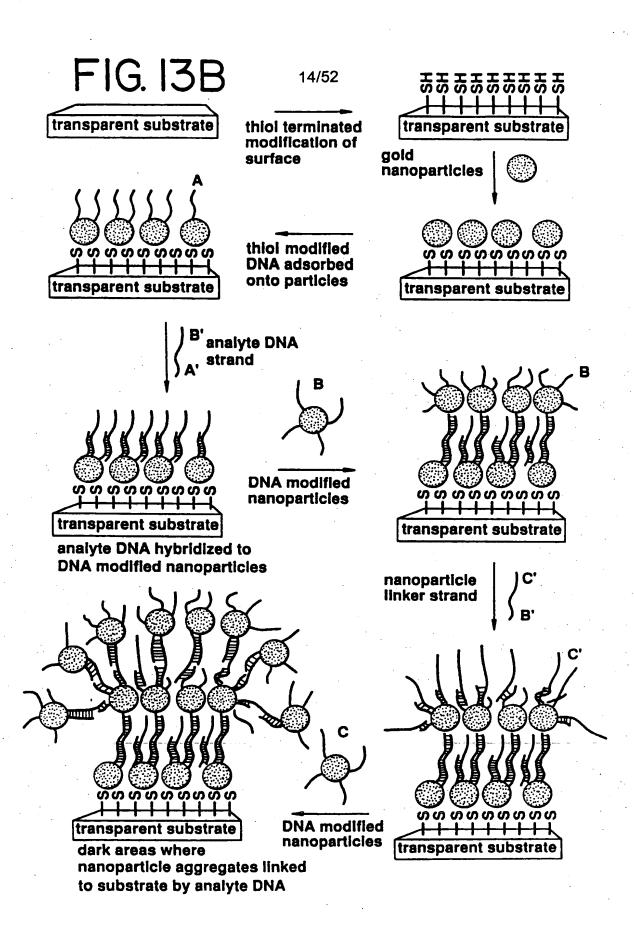
Dark areas where nanoparticle aggregates are linked to substrate surface by analyte DNA

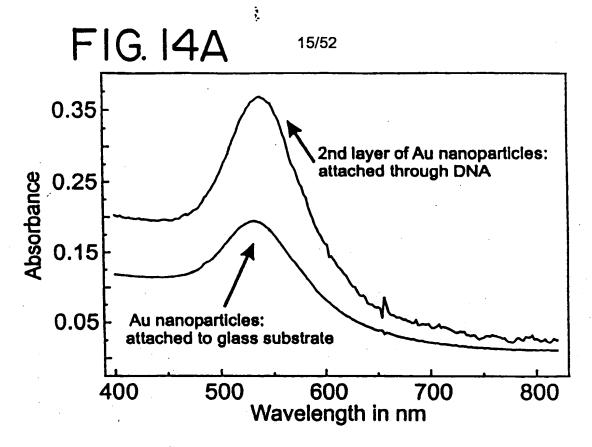




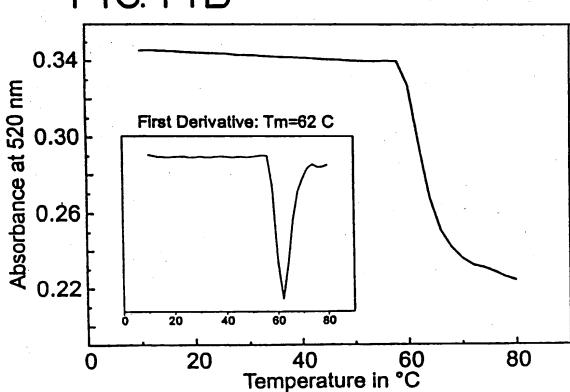
DNA modified colloids

hybridized to bound analyte DNA









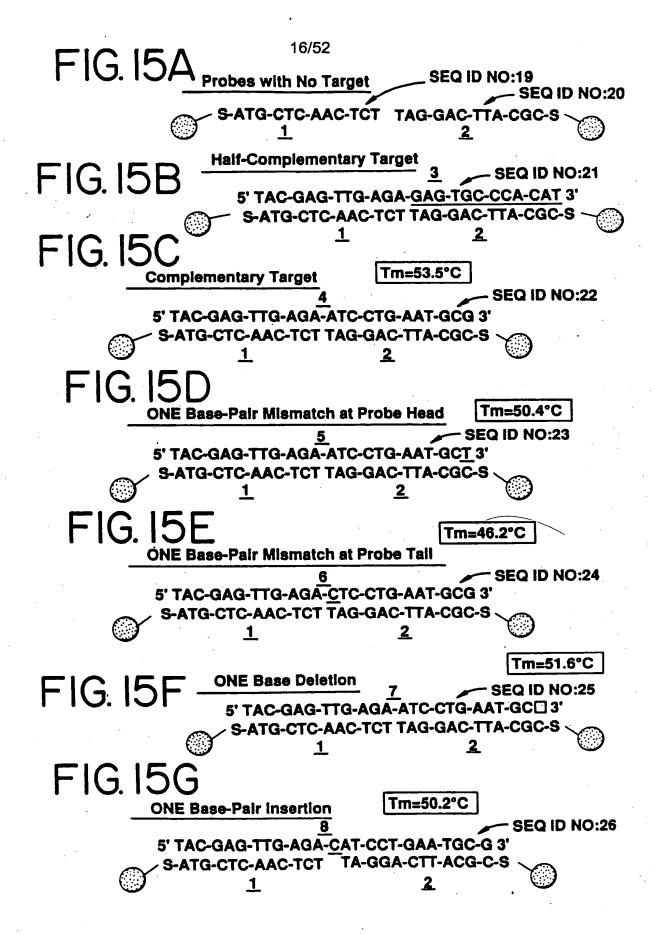


FIG. 16A

24 Base Template

FIG. 16B

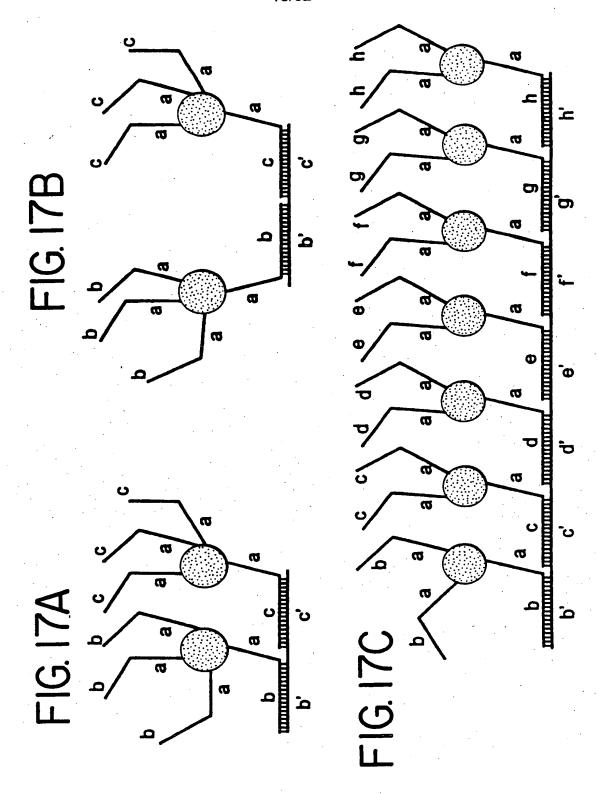
48 Base Template with Complementary 24 Base Filler

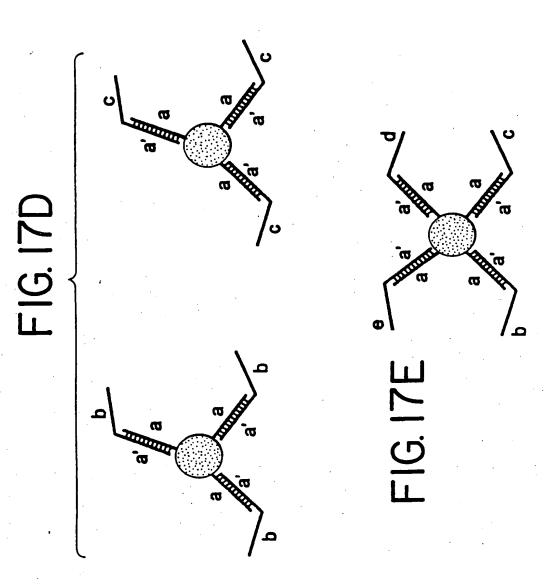
5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-ATC-CTG-AAT-GCG 3' -> S-ATG-CTC-AAC-TCT GGC-AAT-TCT-GCT-CCG-TTA-GTA-CGT TAG-GAC-TTA-CGC-S

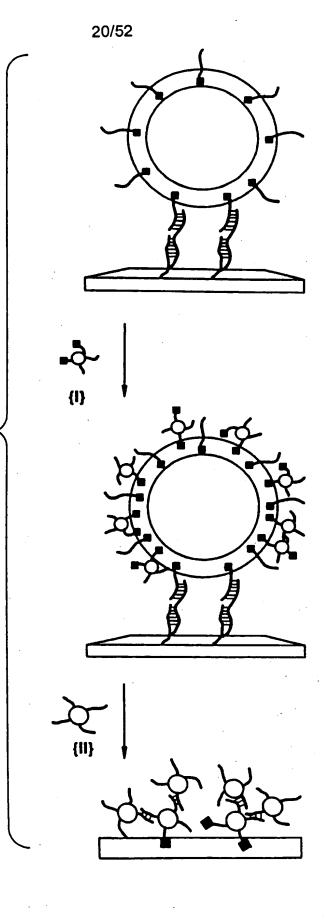
FIG. 16C

72 Base Template with Complementary 48 Base Filler

7







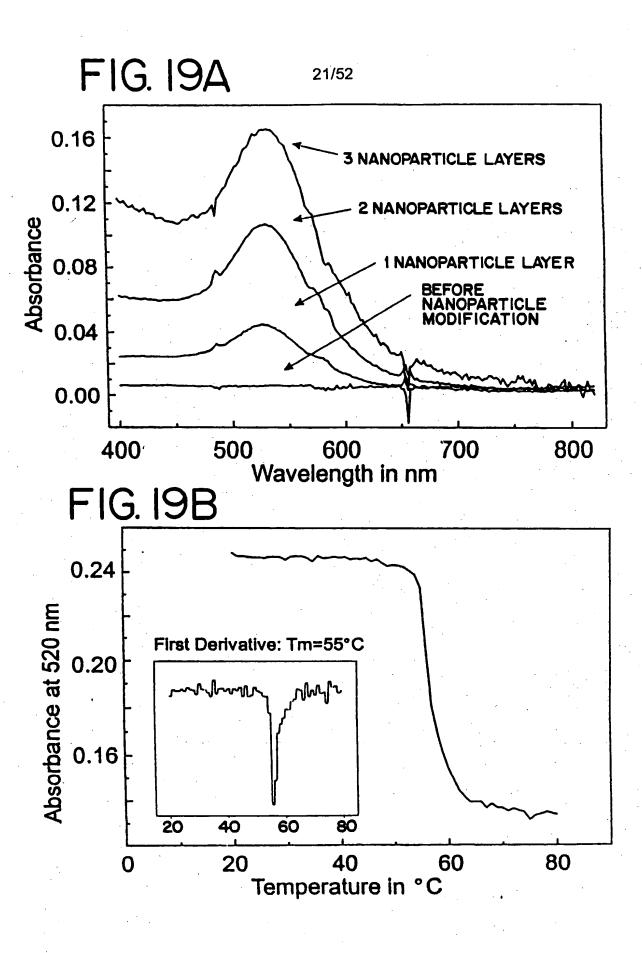
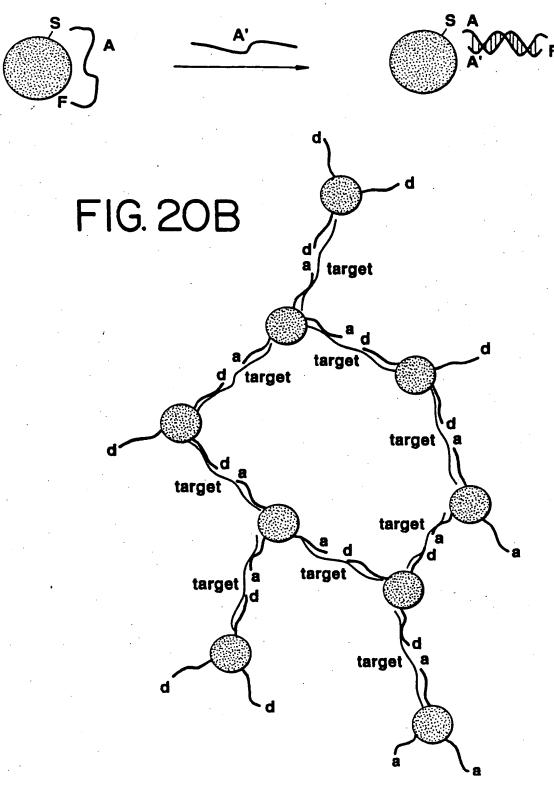
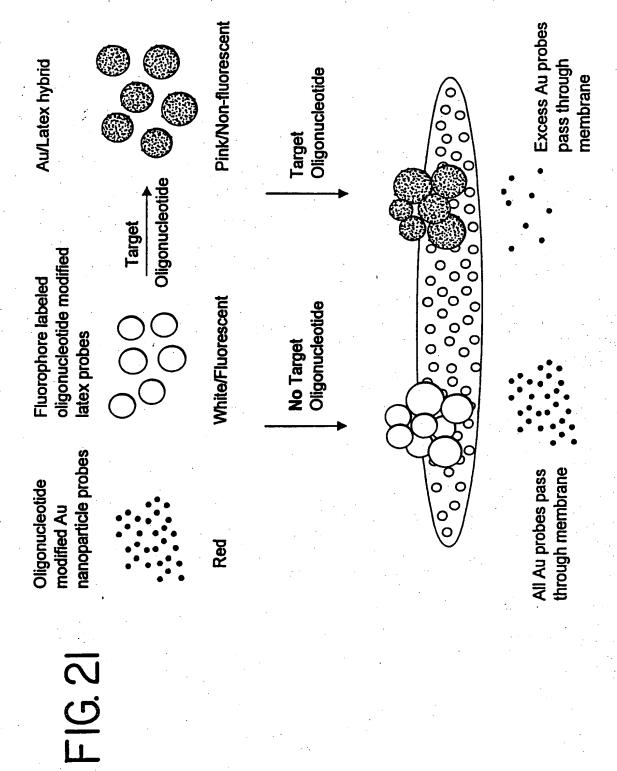
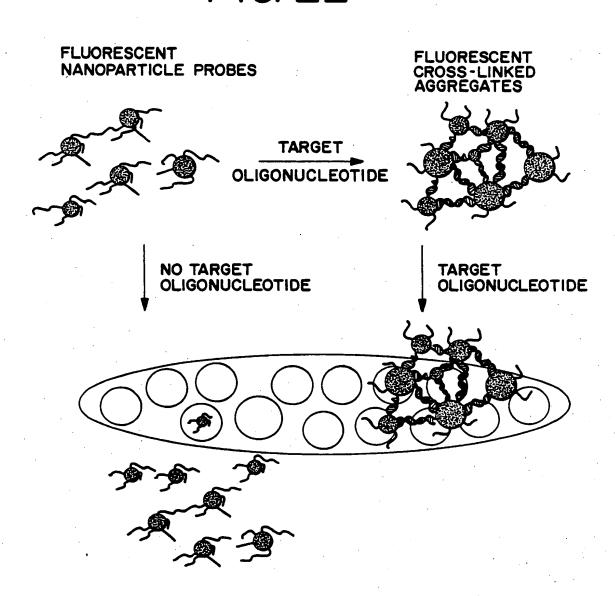


FIG. 20A







THE FLUORESCENT NANOPARTICLE PROBES PASS THROUGH THE MEMBRANE

THE FLUORESCENT CROSS-LINKED AGGREGATES ARE RETAINED BY THE MEMBRANE

Anthrax PCR Product

5'G GCG GAT GAG TCA GTA GTT AAG GAG GCT CAT AGA GAA GTA ATT AAT 3'C CGC CTA CTC AGT CAT CAA TTC CTC CGA GTA TCT CTT CAT TAA TTA

TCG TCA ACA GAG GGA TTA TTG TTA AAT ATT GAT AAG GAT ATA AGA AAA AGC AGT TGT CTC CCT AAT AAC AAT TTA TAA CTA TTC CTA TAT TCT TTT

ATA TTA TCC AGG GTT ATA TTG TAG AAA TTG AAG ATA CTG AAG GGC TT 3'
TAT AAT AGG TCC CAA TAT AAC ATC TTT AAC TTC TAT GAC TTC CCG AA 5'

141 mer Anthrax PCR product [SEQ ID NO:36]

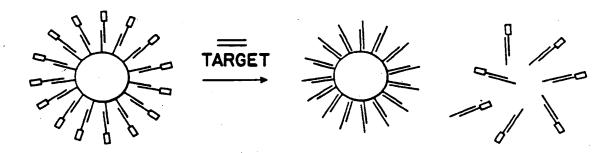
3' CTC CCT AAT AAC AAT

3' TTA TAA CTA TTC CTA (SEQ ID NO:38)

Oligonucleotide-Nanoparticle Probes

Blocker Oligonucleotides

3' C CGC CTA CTC AGT CAT CAA TTC CTC CGA GT	[SEQ ID NO:39]
3' A TCT CTT CAT TAA TTA AGC AGT TGT	[SEQ ID NO:40]
3' TAT TCT TTT TAT AAT AGG TCC CAA TAT	[SEQ ID NO:41]
3' AAC ATC TTT AAC TTC TAT GAC TTC CCG AA	[SEQ ID NO:42]



SATELLITE PROBE

DETECTION SIGNAL

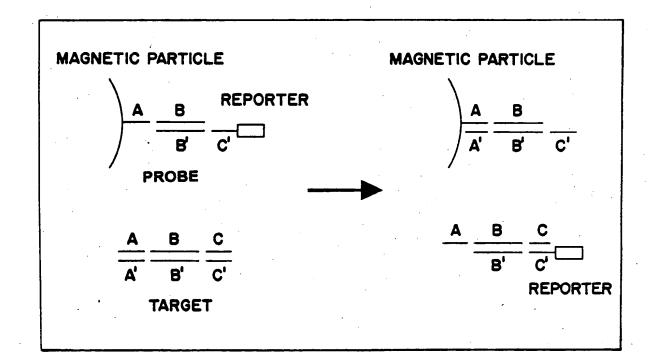
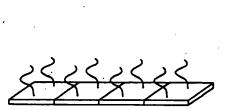


FIG. 25A 1. \sim (target)





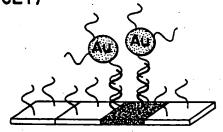
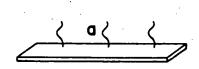
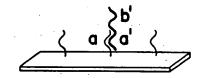
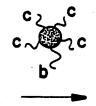


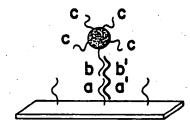
FIG. 25B

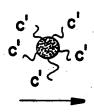


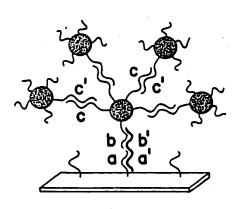












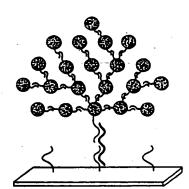


FIG. 26A

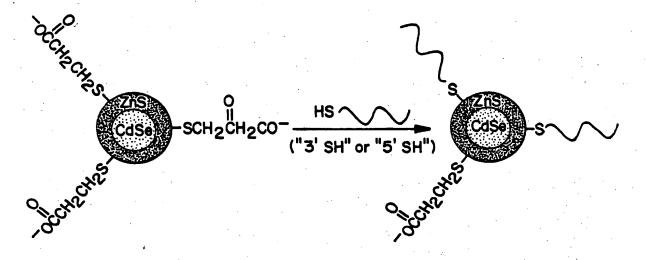


FIG. 26B

TAC-GAG-TTG-AGA-ATC-CTG-AAT-GCG S-[A]_{IO}-ATG-CTC-AAC-TCT TAG-GAC-TTA-CGC-[A]_{IO}S

SEQ ID NO: 46

SEQ ID NO: 47

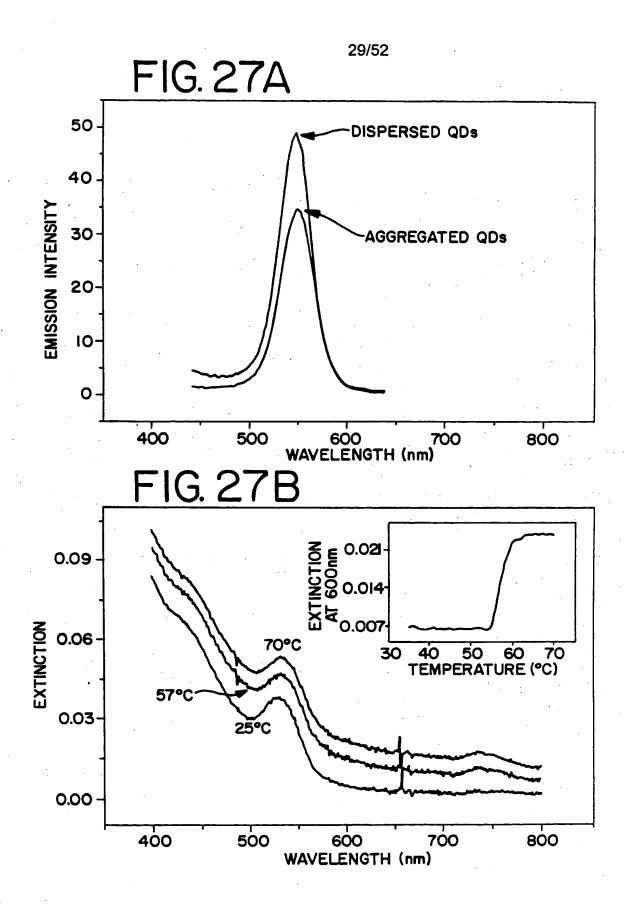


FIG. 27C

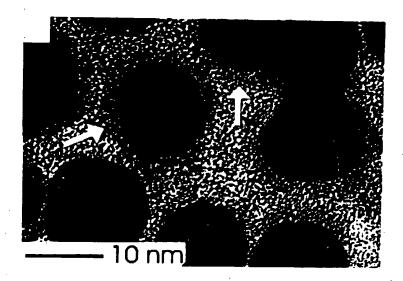


FIG. 27D

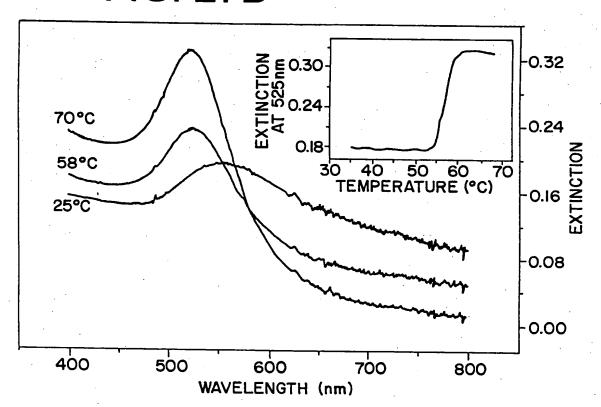


FIG. 28A

FIG. 28B

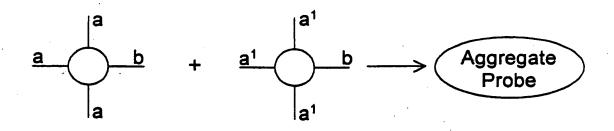


FIG. 28C

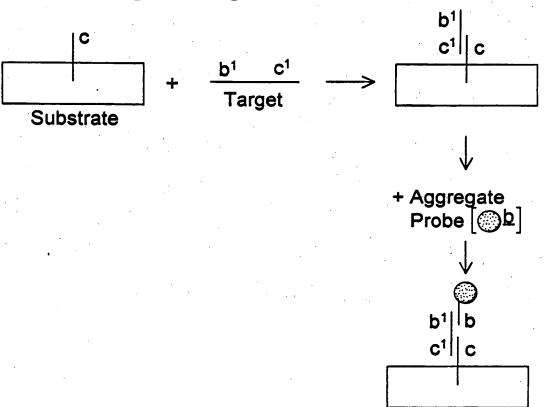


FIG. 28D

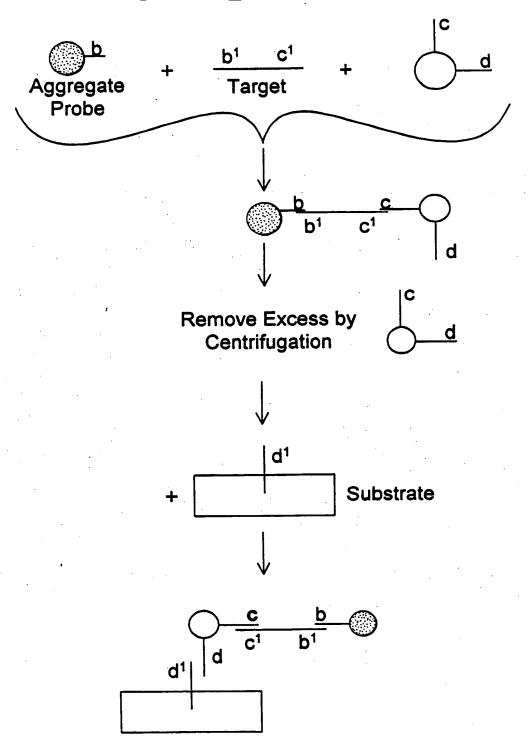
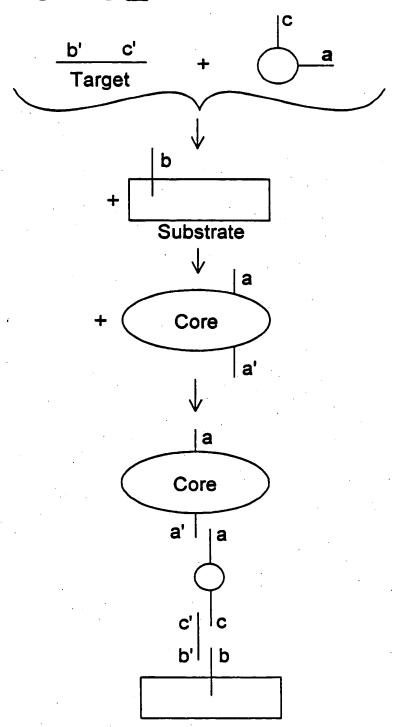
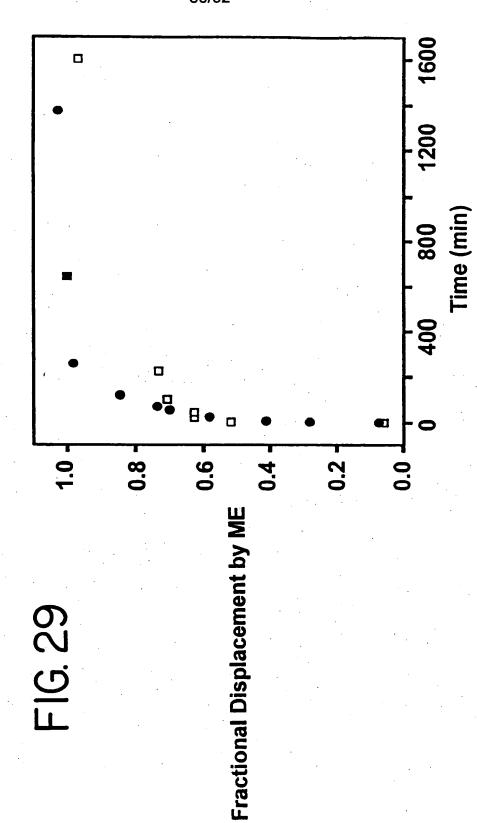
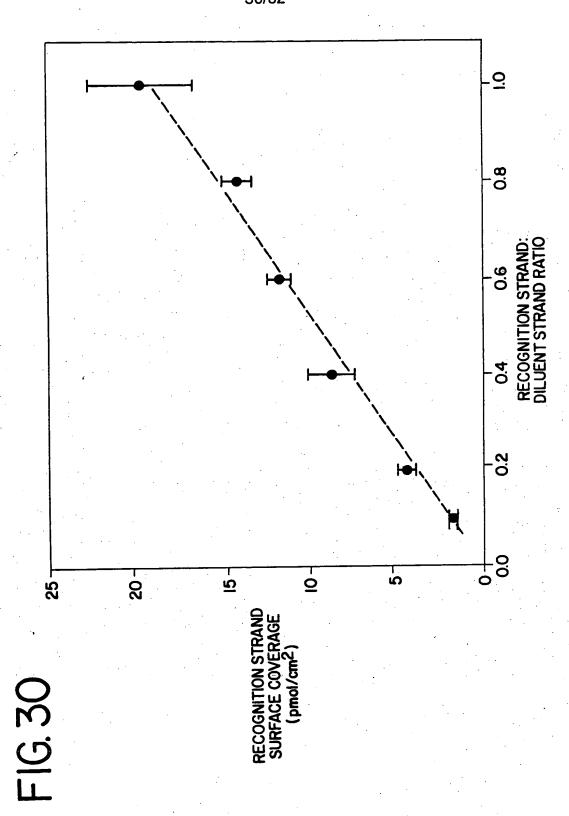
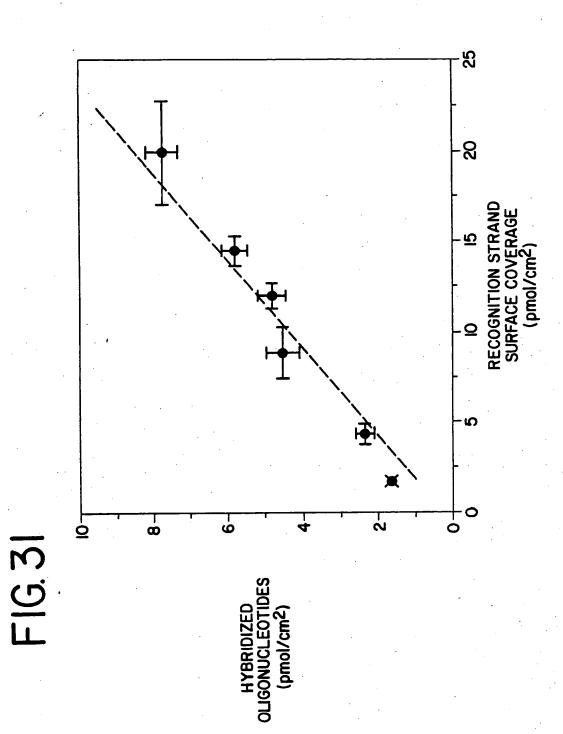


FIG. 28E









[SEQIDNO:56]

5' GGA TTA TTG TTA- -AAT ATT GAT AAG GAT 3'
— CCT ANT AAC AAT TTA TAA CTA TTC CTA

[SEQ ID NO: 57] [SEQ ID NO: 58]

N = A (complementary), G,C,T (mismatched)

1. \((target DNA)

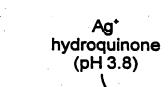
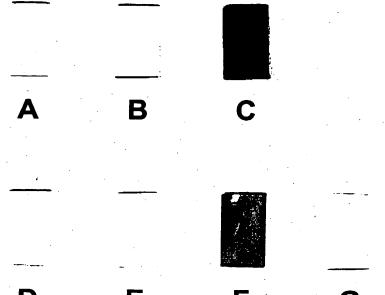
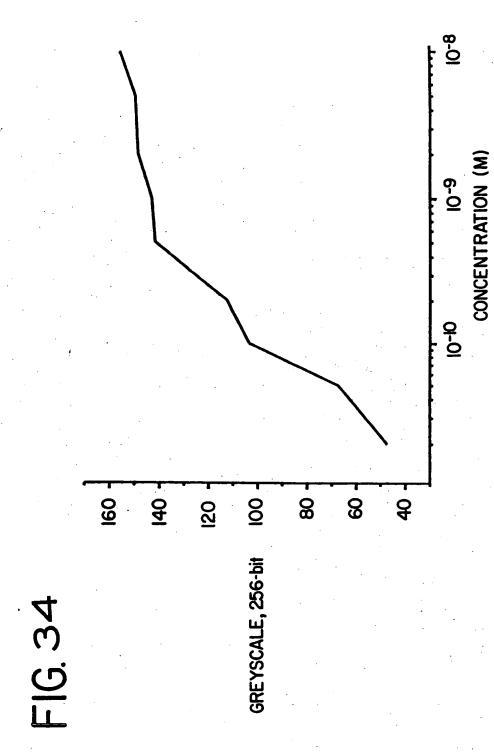


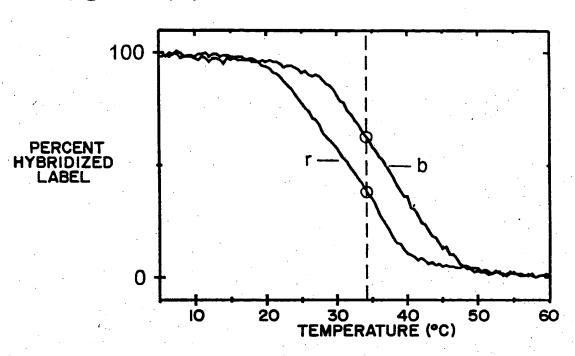


FIG. 33











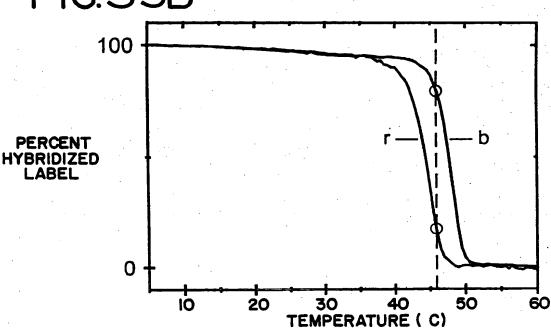


FIG. 36A

FIG. 36B

C A T G

FIG.37A

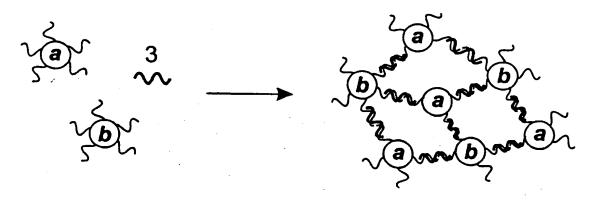
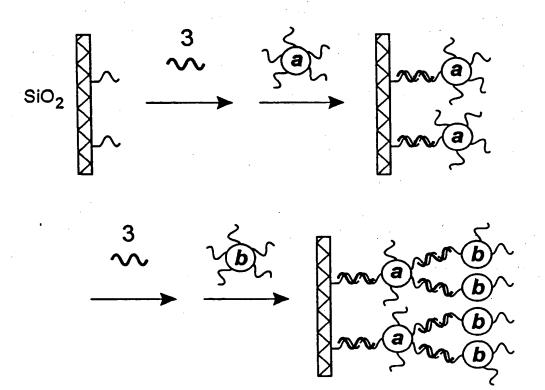
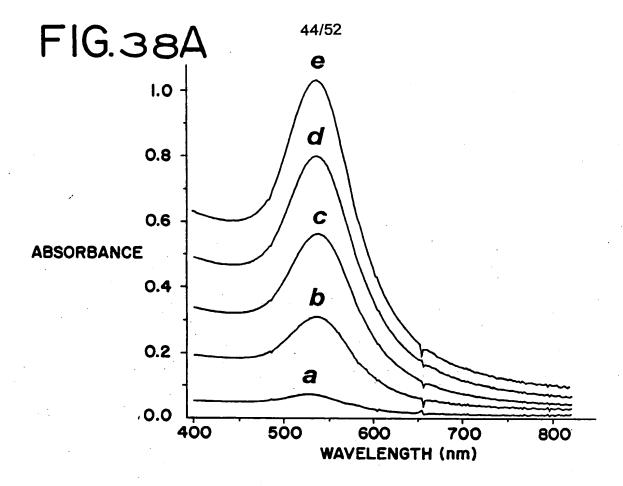


FIG.37B





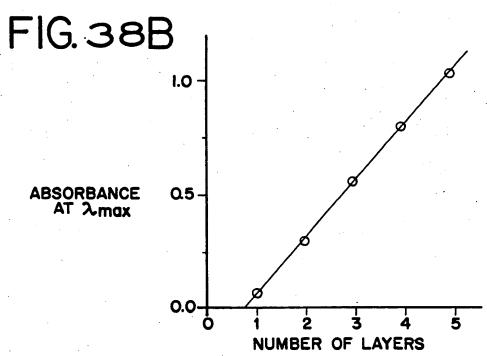
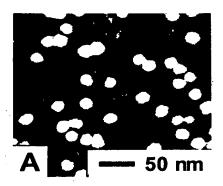
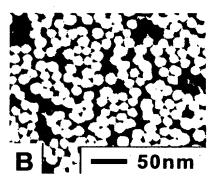
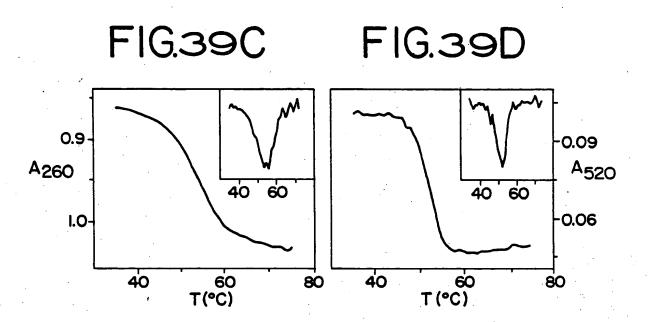


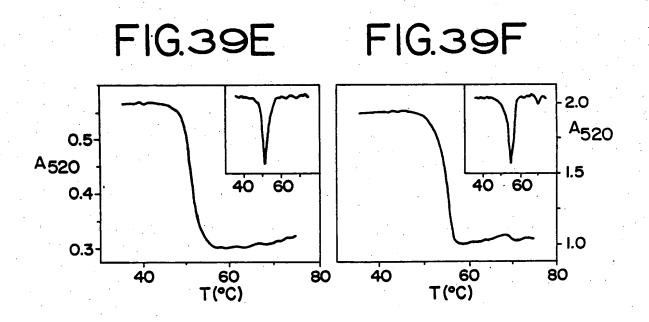
FIG. 39A

FIG. 39B









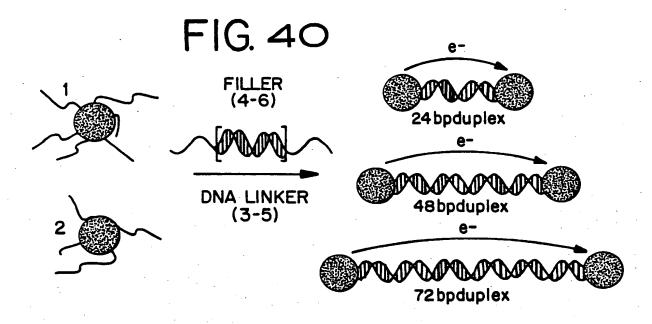
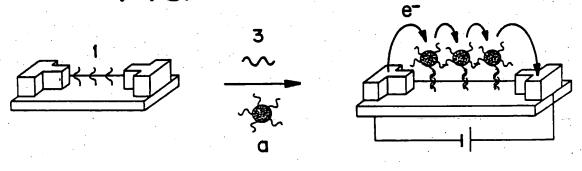


FIG. 41



II HS-(CH₂)₆OR₁

S-(CH₂)₆OR₁ S-(CH₂)₆OH

 R_1

a = H

 $b = (iPr)_2NP(OCH_2CH_2CN)$ -

 $c1 = 5'p(A_{20})$ -TATCGTTCCATCAGCT [SEQ ID NO: 65]

 $c2 = 5'-p(A_{20})-TTGATCTTCCGTTCT$ [SEQ ID NO: 66]

Target I = 79-mer oligonucleotide with target region:

3'------STAGCAAGGTAGTCGAGCAACTAGAAAGGCAAGA......5'
[SEQ ID NO: 67]

49/52 FIG. 43

 R_2 a = H $b = (iPr)_2NP(OCH_2CH_2CN) c1 = 5'-p(A_{20})-GCAGACCTCA \quad [SEQ ID NO: 68]$ $c2 = 5'-p(A_{20})-CCTATGTGTCG \quad [SEQ ID NO: 69]$ $D = 5'-p(A_{20}) \quad [SEQ ID NO: 70]$ Target I = 63-mer oligonucleotide with target region: 3'-....CGTCTGGAGTGGATACACAGC. [SEQ ID NO: 71]

$$R_4$$
-(CH₂)_n R_3 R_4 -NH(CH₂)_n R_3 R_4

R₃ = hydrogen, an alkyl group, an aryl group, or a substituted alkyl or aryl group

 R_4 = an attached oligonucleotide or modified oligonucleotide